



SEQ ID No: 92 DR01006	C-----TCGAG---TCTGGGGGAGG
SEQ ID No: 93 DR27006	C-----TCGAG---TCTGGGGGAGG
SEQ ID No: 94 DR03006	C-----AGGTGA-----AACTGCTCGAG---TCTGGAGGAGG
SEQ ID No: 95 DR11006	C-----TCGAG---TCTGGGGGAGG
SEQ ID No: 96 DR24006	C-----AGGTGA-----AACTGCTCGAG---TCTGGGGGAGG
SEQ ID No: 97 DR16006	C-----TCGAG---TCTGGAGGAGG
SEQ ID No: 98 DR19006	C-----TCGAG---TCTGGAGGAGG
SEQ ID No: 99 DR07006	C-----TCGAG---TCTGGGGGAGG
SEQ ID No: 100 DR16006	C-----TCGAG---TCTGGGGGAGG
SEQ ID No: 101 DR20006	C-----TCGAG---TCAGGGGGAGG
SEQ ID No: 102 DR25006	C-----TCGAG---TCTGGGGGAGG
SEQ ID No: 103 DR20006	C-----TCGAG---TCTGGAGGAGG
SEQ ID No: 104 DR21006	C-----TCGAG---TCTGGGGGAGG
SEQ ID No: 105 DR09006	C-----AGGTGA-----AACTGCTCGAG---TCTGGGGGAGG
SEQ ID No: 106 DR17006	C-----TCGAG---TCTGGGGGAGG
SEQ ID No: 107 DR13006	C-----TCGAG---TCAGGGGGAGG
SEQ ID No: 108 DR02006	CTCGAGTCAGGTGTCCGGTCTGATGTGCAGCTGGTGGCGTCTGGGGGAGG
DR01006	ATCGGTGCAGGCTGGAGGGTCTCTGAGACTCTC--GTGCG-CAGCCTCTG
DR27006	CTCGGTGCAGGCTGGAGGGTCTCTGAGACTCTCCTGTGCATCTTCTTCTA
DR03006	CTCGGTGCAGACTGGAGGATCTCTGAGACTCTCCTGTGCAGT--C-TCTG
DR11006	GTCGGTGCAGGCTGGAGGGTCTCTGAGACTCTCCTGTAATGT--C-TCTG
DR24006	GTCGGTGCAGGCTGGAGGGTCTCTGAGACTCTCCTGTAATGT--C-TCTG
DR16006	CTCGGCGCAGGCTGGAGGATCTCTGAGACTCTCCTGTGCAGC--CCACGG
DR19006	CTCGGTTTCAGGCTGGAGGGTCCCTTAGACTCTCCTGTGCAGC--C-TCTG
DR07006	CTCGGTGCAGGGTGGAGGGTCTCTGAGACTCTCCTGTGCAA---TCTCTG
DR16006	CTCGGTGCAGGCTGGAGGGTCTCTGAGACTCTCCTGTACAG---GCTCTG
DR20006	CTCGGTACAGGTTGGAGGGTCTCTGAGACTCTCCTGTGTAG---CCTCTA
DR25006	CTCGGTACAAACTGGAGGGTCTCTGAGACTCTCCTGTGCG---AAATCTCTG
DR20006	CTCGGTGCAGGCTGGAGGGTCTCTGAGACTCTCCTGTG---TAGCCTCTG
DR21006	CTCGGTGCAGGTTGGAGGGTCTCTGAAACTCTCCTGTAAAAT---CTCTG
DR09006	CTCGGTGCAGGCTGGGGGTCTCTGACACTCTCCTGTAGTT---TATACAC--
DR17006	CTCGGTCCAACCTGGAGGATCTCTGACACTCTCCTGTACAGTT---TCTG
DR13006	CTCGGTGGAGGCTGGAGGGTCTCTGAGACTCTCCTGTACAG---CCTCTG
DR02006	CTCGGTGCAGGCTGGAGGCTCTCTGAGACTCTCCTGTACAG---CCTCTG
DR01006	GA--TACAGTAATT---GTCCCCTCACTTG-GAGCTGGTATCGCCAGTTT
DR27006	AA--TATATGCCTT---GCACCTACGACAT-GACCTGGTACCGCCAGGCT
DR03006	GA--TTCTCCTTTA---GTACCAGTTGTAT-GGCCTGGTTCCGCCAGGCT
DR11006	GC--TCTCCCAGTA---GTACTTATTGCCT-GGGCTGGTTCCGCCAGGCT
DR24006	GC--TCTCCCAGTA---GTACTTATTGCCT-GGGCTGGTTCCGCCAGGCT
DR16006	GA--TTCCGC-TCA---ATGGTTACTACAT-CGCCTGGTTCCGTCAGGCT
DR19006	AC--TACACCATCA---CTGATTATTGCAT-GGCCTGGTTCCGCCAGGCT
DR07006	GA--TACACGTACG---GTAGCTTCTGTAT-GGGCTGGTTCCGCGAGGGT
DR16006	GA--TTCCCCTATA---GTACCTTCTGTCT-GGGGTGGTTCCGCCAGGCT
DR20006	CT--CACACCGACA---GTAGCACCTGTAT-AGGCTGGTTCCGCCAGGCT
DR25006	GA--TTGACTTTTG---ATGATTCTGACGT-GGGGTGGTACCGCCAGGCT
DR20006	GA--TTCAATTTTG---AACTTCTCGTAT-GGCCTGGTACCGCCAGACT
DR21006	GAGGTACCCAGATCGTGTTCCTAAATCTTTGGCCTGGTTCCGCCAGGCT
DR09006	-----CAACGATACTGGGACCA-----TGGGATGGTTTCGCCAGGCT
DR17006	--GGGCCACCTACA---GTGACTACAGTATTGGA-TGGATCCGCCAGGCT
DR13006	G-----ATACGTAT-CCT----CTATGGCCTGGTTCCGCCAGGTT
DR02006	GAGA----CAGTTTCAGTAGATT--TGCCATGTCTTGGTTCCGCCAGGCT

FIG. 7A